

Sequence Homology Search

RESULT 1

US-09-782-906-2

; Sequence 2, Application US/09782906

; Patent No. US20010051369A1

; GENERAL INFORMATION:

; APPLICANT: Delagrave, Simon

; APPLICANT: Rittenhouse Pruss, Jennifer L.

; APPLICANT: Murphy, Dennis J.

; APPLICANT: Maffia III, Anthony M.

; APPLICANT: Bylina, Edward J.

; APPLICANT: Coleman, William J.

; TITLE OF INVENTION: Variant Galactose Oxidase, Nucleic Acid Encoding Same,
And Methods Of

; TITLE OF INVENTION: Using Same

; FILE REFERENCE: HER-0040

; CURRENT APPLICATION NUMBER: US/09/782,906

; CURRENT FILING DATE: 2001-02-14

; PRIOR APPLICATION NUMBER: 60/185,001

; PRIOR FILING DATE: 2000-02-25

; NUMBER OF SEQ ID NOS: 8

; SOFTWARE: PatentIn version 3.0

; SEQ ID NO 2

; LENGTH: 639

; TYPE: PRT

; ORGANISM: Dactylium dendroides

US-09-782-906-2

Query Match 94.4%; Score 3425; DB 10; Length 639;

Best Local Similarity 100.0%; Pred. No. 4.2e-255;

Matches 639; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy      42 ASAPIGSAISRNNWAVTCDSAQSGNECNKAIDGNKDTFWHTFYGANGDPKPPHTYTTIDMK 101
         |||
Db      1 ASAPIGSAISRNNWAVTCDSAQSGNECNKAIDGNKDTFWHTFYGANGDPKPPHTYTTIDMK 60

Qy     102 TTQNVNGLSMLPRQDGNQNGWIGRHEVYLSSDGTNWGSPVASGSWFADSTTKYSNFETRP 161
         |||
Db      61 TTQNVNGLSMLPRQDGNQNGWIGRHEVYLSSDGTNWGSPVASGSWFADSTTKYSNFETRP 120

Qy     162 ARYVRLVAITEANGQPWTSIAEINVQASSYTAPQPGLGRWGPTIDLPIVAAAAIEPTS 221
         |||
Db     121 ARYVRLVAITEANGQPWTSIAEINVQASSYTAPQPGLGRWGPTIDLPIVAAAAIEPTS 180

Qy     222 GRVLMWSSYRNDAFGGSPGGITLTSSWDPSTGIVSDRTVTVTKHMFCPGISMDGNGQIV 281
         |||
Db     181 GRVLMWSSYRNDAFGGSPGGITLTSSWDPSTGIVSDRTVTVTKHMFCPGISMDGNGQIV 240

Qy     282 VTGGNDAKKTSLYDSSSDSWIPGPDMPQVARGYQSSATMSDGRVFTIGGSWSGGVFEKNGE 341
         |||
Db     241 VTGGNDAKKTSLYDSSSDSWIPGPDMPQVARGYQSSATMSDGRVFTIGGSWSGGVFEKNGE 300

Qy     342 VYSPSSKTTWTSLPNAKVNPMILTADKQGLYRSDNHAWLFGWKKGSVFQAGPSTAMNYYTS 401
         |||
Db     301 VYSPSSKTTWTSLPNAKVNPMILTADKQGLYRSDNHAWLFGWKKGSVFQAGPSTAMNYYTS 360

Qy     402 GSGDVKSAGKRQSNRGVAPDAMCGNAVMYDAVKGKILTFGGSPDYQSDATTNNAHIITLG 461
         |||
Db     361 GSGDVKSAGKRQSNRGVAPDAMCGNAVMYDAVKGKILTFGGSPDYQSDATTNNAHIITLG 420

Qy     462 EPGTSPNTVFASNGLYFARTFHTSVVLPDGGSTFITGGQRRGIPFEDSTPVFTPEIYVPEQ 521
         |||
Db     421 EPGTSPNTVFASNGLYFARTFHTSVVLPDGGSTFITGGQRRGIPFEDSTPVFTPEIYVPEQ 480

Qy     522 DTFYKQNPNSIVRVYHSISLLLPDGRVFNNGGGGLCGDCTTNHFDAQIFTNPNLYNSNGNL 581
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Db     481 DTFYKQNPNSIVRVYHSISLLLPDGRVFNNGGGGLCGDCTTNHFDAQIFTNPNLYNSNGNL 540

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Qy      582 ATRPKITRTSTQSVKVGGRITISTDSSISKASLIRYGTATHTVNTDQRRIPLTLTNNGGN 641
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Db      541 ATRPKITRTSTQSVKVGGRITISTDSSISKASLIRYGTATHTVNTDQRRIPLTLTNNGGN 600
        |||

Qy      642 SYSFQVPSDSGVALPGYWMLFVMNSAGVPSVASTIRVTQ 680
        |||
Db      601 SYSFQVPSDSGVALPGYWMLFVMNSAGVPSVASTIRVTQ 639

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Qy      43 ASAPIGSAISRNNWAVTCDSAQSGNECNKAIDGNKDTFWHTFYGANGDPKPPHTYTIDMK 102
         |||
Db       1 ASAPIGSAISRNNWAVTCDSAQSGNECNKAIDGNKDTFWHTFYGANGDPKPPHTYTIDMK 60

Qy     103 TTQNVNGLSMLPRQDGNQNGWIGRHEVYLSSDGTNWGSPVASGSWFADSTTKYSNFETRP 162
         |||
Db       61 TTQNVNGLSMLPRQDGNQNGWIGRHEVYLSSDGTNWGSPVASGSWFADSTTKYSNFETRP 120

Qy     163 ARYVRLVAITEANGQPWTSIAEINVQASSYTAPQPGLGRWGPTIDLPIVPA AAAIEPTS 222
         |||
Db      121 ARYVRLVAITEANGQPWTSIAEINVQASSYTAPQPGLGRWGPTIDLPIVPA AAAIEPTS 180

Qy     223 GRVLMWSSYRNDAFGGSPGGITLTSSWDPSTGIVSDRTVTVTKHMFCPGISMDGNGQIV 282
         |||
Db      181 GRVLMWSSYRNDAFGGSPGGITLTSSWDPSTGIVSDRTVTVTKHMFCPGISMDGNGQIV 240

Qy     283 VTGGNDAKKTSLYDSSSDSWIPGPD MQVARGYQSSATMSDGRVFTIGGSWSGGVFEKNGE 342
         |||
Db      241 VTGGNDAKKTSLYDSSSDSWIPGPD MQVARGYQSSATMSDGRVFTIGGSWSGGVFEKNGE 300

Qy     343 VYSPSSKTTWTS L PNAKVNPMLTADKQGLYRSDNHAWLFGWKKGSVFQAGPSTAMN WYYTS 402
         |||
Db      301 VYSPSSKTTWTS L PNAKVNPMLTADKQGLYRSDNHAWLFGWKKGSVFQAGPSTAMN WYYTS 360

Qy     403 GSGDVKSAGKRQSNRGVAPDAMCGNAV MYDAVKGKILTFGGSPDYQSDATTN AHIITLG 462
         |||
Db      361 GSGDVKSAGKRQSNRGVAPDAMCGNAV MYDAVKGKILTFGGSPDYQSDATTN AHIITLG 420

Qy     463 EPGTSPNTVFASNGLYFARTFHTSVVLPD GSTFITGGQRRGIPFEDSTPVFTPEIYVPEQ 522
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Db      421 EPGTSPNTVFASNGLYFARTFHTSVVLPD GSTFITGGQRRGIPFEDSTPVFTPEIYVPEQ 480

Qy     523 DTFYKQNPNSIVRVYHSIS LLLPDGRVFN GGGGLCGDCTTNHFDAQIFT PNYLYNSNGNL 582
         |||
Db      481 DTFYKQNPNSIVRVYHSIS LLLPDGRVFN GGGGLCGDCTTNHFDAQIFT PNYLYNSNGNL 540

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Qy 583 ATRPKITRTSTQSVKVGGRITISTDSSISKASLIRYGTATHTVNTDQRRIPLTLTNNGN 642
|||||
Db 541 ATRPKITRTSTQSVKVGGRITISTDSSISKASLIRYGTATHTVNTDQRRIPLTLTNNGN 600
Qy 643 SYSFQVPSDSGVALPGYWMLFVMNSAGVPSVASTIRVTQ 681
|||||
Db 601 SYSFQVPSDSGVALPGYWMLFVMNSAGVPSVASTIRVTQ 639